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Supplementary appendix

This appendix formed part of the original submission and has been peer reviewed. We post it as supplied by the authors.

This online publication has been corrected. The corrected version first appeared at thelancet.com on August 20, 2020.

Supplement to: Lee LYW, Cazier JB, Angelis V, et al. COVID-19 mortality in patients with cancer on chemotherapy or other anticancer treatments: a prospective cohort study. *Lancet* 2020; published online May 28. https://doi.org/10.1016/S0140-6736(20)31173-9.

Supplementary Methods

Data visualisation and figure generation

Data processing and visualisation utilised R (version 3.6.3) packages including broom, dplyr, gpclib, ggmap, ggplot2, mapdata, maps, maptools, networkD3, rgdal, rgeos, robustbase and viridis. Data subsetting was performed using the subset() function of 'robustbase' and data reshaping for visualisation involved the use of the group_by() and melt() functions of 'dplyr'. Functions from the ggplot2 R package were used to generate multiple plots including barplots (geom_bar) and UK region map (geom_polygon). The sankeyNetwork() function of the 'networkD3' R package was also used to generate the Sankey plot. The shape (.shp) file for the UK region map was publicly available from the UK Office for National Statistics.

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